AASS gene

aminoadipate-semialdehyde synthase

Normal Function

The AASS gene provides instructions for making an enzyme called aminoadipic semialdehyde synthase. This enzyme is found in most tissues, with the highest amounts found in the liver. Aminoadipic semialdehyde synthase is involved in the breakdown of the amino acid lysine, a building block of most proteins. It is called a bifunctional enzyme because is performs two functions. One function, called lysine-ketoglutarate reductase, breaks down lysine to a molecule called saccharopine. The other function, called saccharopine dehydrogenase, breaks down saccharopine to a molecule called alpha-aminoadipate semialdehyde.

Health Conditions Related to Genetic Changes

<u>hyperlysinemia</u>

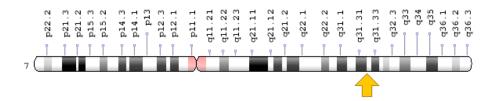
At least five mutations in the AASS gene have been found to cause hyperlysinemia. Most of these mutations change single amino acids in aminoadipic semialdehyde synthase. These mutations are thought to decrease or eliminate enzyme activity, resulting in an inability to break down lysine. Lysine that is not broken down accumulates in the blood, but it typically causes no health problems.

When mutations in the *AASS* gene impair the breakdown of saccharopine, this molecule builds up in blood and urine. This buildup is sometimes referred to as saccharopinuria, which is considered to be a variant of hyperlysinemia. It is unclear if saccharopinuria causes any symptoms.

Chromosomal Location

Cytogenetic Location: 7q31.32, which is the long (q) arm of chromosome 7 at position 31.32

Molecular Location: base pairs 122,073,544 to 122,144,290 on chromosome 7 (Homo sapiens Annotation Release 108, GRCh38.p7) (NCBI)



Credit: Genome Decoration Page/NCBI

Other Names for This Gene

- AASS_HUMAN
- alpha-aminoadipate semialdehyde synthase
- aminoadipic semialdehyde synthase
- LKR/SDH
- LKRSDH
- lysine-2-oxoglutarate reductase
- lysine-ketoglutarate reductase /saccharopine dehydrogenase

Additional Information & Resources

Genetic Testing Registry

 GTR: Genetic tests for AASS https://www.ncbi.nlm.nih.gov/gtr/all/tests/?term=10157%5Bgeneid%5D

Scientific Articles on PubMed

PubMed

https://www.ncbi.nlm.nih.gov/pubmed?term=%28%28AASS%5BTIAB%5D%29+OR+%28aminoadipate-semialdehyde+synthase%29%29+AND+%28%28Genes%5BMH%5D%29+OR+%28Genetic+Phenomena%5BMH%5D%29%29+AND+english%5Bla%5D+AND+human%5Bmh%5D+AND+%22last+3600+days%22%5Bdp%5D

OMIM

 ALPHA-AMINOADIPIC SEMIALDEHYDE SYNTHASE http://omim.org/entry/605113

Research Resources

- ClinVar https://www.ncbi.nlm.nih.gov/clinvar?term=AASS%5Bgene%5D
- HGNC Gene Symbol Report http://www.genenames.org/cgi-bin/gene_symbol_report?q=data/ hgnc_data.php&hgnc_id=17366
- NCBI Gene https://www.ncbi.nlm.nih.gov/gene/10157
- UniProt http://www.uniprot.org/uniprot/Q9UDR5

Sources for This Summary

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- Sacksteder KA, Biery BJ, Morrell JC, Goodman BK, Geisbrecht BV, Cox RP, Gould SJ, Geraghty MT. Identification of the alpha-aminoadipic semialdehyde synthase gene, which is defective in familial hyperlysinemia. Am J Hum Genet. 2000 Jun;66(6):1736-43. Epub 2000 Apr 20. Citation on PubMed: https://www.ncbi.nlm.nih.gov/pubmed/10775527
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